SEQUENCE LISTING

<110> HITACHI,LTD

<120> Method for Testing Nucleic Acid Sequence

<130> WO216-01CW; ASAM.0055

<140> US 10/083,340
<141> 2002-02-27

<150> JP 2001-331853
<151> 2001-10-30

<160> 4

<210> 1
<211> 95
<212> DNA
<213> Artificial Sequence

<220>
<223> Sequence of portion of model gene including mutation.

<400> 1
ctttcttgcg gagattctct tcctctgtgc gccggtctct cccaggacag gcacaaacac 60
gcacctcaaa gctgttccgt cccagtagat tacca   95

<210> 2
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA probe for detecting wild type and having an artificial
      mismatched base at the third base from the 3'end.

<400> 2
aacagctttg aggtgcgtga ta 22

<210> 3
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> DNA probe for detecting mutation type and having an artificial
      mismatched base at the third base from the 3'end.

<400> 3
aacagctttg aggtgcgtga tt 22

<210> 4
<211> 22
<212> DNA
<213> Artificial Sequence

<220>
<223> Sequence portion of SEQ.No.1 which is a target sequence
      hybridized with DNA probe having base sequence of SEQ.No.2 or

No.3.

<400> 4
aaacacgcac ctcaaagctg tt 22

INPUT SEQ: A:\ASAM55.ans

GENERAL INFORMATION SECTION

    3,<110> HITACHI,LTD.
    5,<120> Method for Testing Nucleic Acid Sequence
    7,<130> WO216-01CW; ASAM.0055
    9,<140> US 10/083,340
  10,<141> 2002-02-27
  12,<150> JP 2001-331853
  13,<151> 2001-10-30
  15,<160> 4

ERRORED LINES SECTION

STATISTICS SUMMARY

    Application Serial Number: US 10/083,340
        Alpha or Numeric: Numeric
       Application Class:
    Application File Date: 2002-02-27
           Art Unit:
    Software Application:
Total Number of Sequences: 4
     Total Nucleotides: 161
     Total Amino Acids: 0
     Number of Errors: 0
   Number of Warnings: 0
Number of Corrections: 0